

SEQUENCE LISTING

<110> Higuchi, Maria de Lourdes
Schenkman, Sergio

<120> PREVENTION AND TREATMENT OF
MYCOPLASMA-ASSOCIATED DISEASES

<130> 33474-PCT-USA-A 068528.0103

<140> Not Yet Assigned

<141> 2002-01-03

<150> PCT/BR01/00083

<151> 2001-03-07

<150> Not Yet Assigned

<151> 2001-03-07

<150> PI 0002989-0 BR

<151> 2000-03-07

<160> 4

<170> FastSEQ for Windows Version 4.0

<210> 1

<211> 2010

<212> DNA

<213> Artificial Sequence

<220>

<223> Variant of T. Cruzi trans-sialidase gene

<400> 1

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<210> 2

<211> 669

<212> PRT

<213> Artificial Sequence

<220>

<223> Variant of T. Cruzi trans-sialidase protein

<400> 2

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          35          40          45
Arg Val Val His Ser Phe Arg Leu Pro Ala Leu Val Asn Val Asp Gly
          50          55          60
Val Met Val Ala Ile Ala Asp Ala Arg Tyr Glu Thr Ser Asn Asp Asn
65          70          75          80
Ser Leu Ile Asp Thr Val Ala Lys Tyr Ser Val Asp Asp Gly Glu Thr
          85          90          95
Trp Glu Thr Gln Ile Ala Ile Lys Asn Ser Arg Ala Ser Ser Val Ser
          100          105          110
Arg Val Val Asp Pro Thr Val Ile Val Lys Gly Asn Lys Leu Tyr Val
          115          120          125
Leu Val Gly Ser Tyr Asn Ser Ser Arg Ser Tyr Trp Thr Ser His Gly
          130          135          140
Asp Ala Arg Asp Trp Asp Ile Leu Leu Ala Val Gly Glu Val Thr Lys
145          150          155          160
Ser Thr Ala Gly Gly Lys Ile Thr Ala Ser Ile Lys Trp Gly Ser Pro
          165          170          175
Val Ser Leu Lys Glu Phe Phe Pro Ala Glu Met Glu Gly Met His Thr
          180          185          190
Asn Gln Phe Leu Gly Gly Ala Gly Val Ala Ile Val Ala Ser Asn Gly
          195          200          205
Asn Leu Val Tyr Pro Val Gln Val Thr Asn Lys Lys Lys Gln Val Phe
210          215          220
Ser Lys Ile Phe Tyr Ser Glu Asp Glu Gly Lys Thr Trp Lys Phe Gly
225          230          235          240
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Glu Gly Lys Leu Ile Ile Asn Thr Arg Val Asp Tyr Arg Arg Arg Leu
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Thr	Leu	Ser	Arg	Val	Trp	Gly	Pro	Ser	Pro	Lys	Ser	Asn	Gln	Pro	Gly	290	295	300
Ser	Gln	Ser	Ser	Phe	Thr	Ala	Val	Thr	Ile	Glu	Gly	Met	Arg	Val	Met	305	310	315
Leu	Phe	Thr	His	Pro	Leu	Asn	Phe	Lys	Gly	Arg	Trp	Leu	Arg	Asp	Arg	320	325	330
Leu	Asn	Leu	Trp	Leu	Thr	Asp	Asn	Gln	Arg	Ile	Tyr	Asn	Val	Gly	Gln	335	340	345
Val	Ser	Ile	Gly	Asp	Glu	Asn	Ser	Ala	Tyr	Ser	Ser	Val	Leu	Tyr	Lys	350	355	360
Asp	Asp	Lys	Leu	Tyr	Cys	Leu	His	Glu	Ile	Asn	Ser	Asn	Glu	Val	Tyr	365	370	375
Ser	Leu	Val	Phe	Ala	Arg	Leu	Val	Gly	Glu	Leu	Arg	Ile	Ile	Lys	Ser	380	385	390
Val	Leu	Gln	Ser	Trp	Lys	Asn	Trp	Asp	Ser	His	Leu	Ser	Ser	Ile	Cys	395	400	405
Thr	Pro	Ala	Asp	Pro	Ala	Ala	Ser	Ser	Ser	Glu	Arg	Gly	Cys	Gly	Pro	410	415	420
Ala	Val	Thr	Thr	Val	Gly	Leu	Val	Gly	Phe	Leu	Ser	His	Ser	Ala	Thr	425	430	435
Lys	Thr	Glu	Trp	Glu	Asp	Ala	Tyr	Arg	Cys	Val	Asn	Ala	Ser	Thr	Ala	440	445	450
Asn	Ala	Glu	Arg	Val	Pro	Asn	Gly	Leu	Lys	Phe	Ala	Gly	Val	Gly	Gly	455	460	465
Gly	Ala	Leu	Trp	Pro	Val	Ser	Gln	Gln	Gly	Gln	Asn	Gln	Arg	Tyr	His	470	475	480
Phe	Ala	Asn	His	Ala	Phe	Thr	Leu	Val	Ala	Ser	Val	Thr	Ile	His	Glu	485	490	495
Val	Pro	Ser	Val	Ala	Ser	Pro	Leu	Gly	Ala	Ser	Leu	Asp	Ser	Ser		500	505	510
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<210> 3

<211> 28

<212> DNA

<213> Artificial Sequence

<223> Trans-sialidase gene primer

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28

<211> 34

<212> DNA

<213> Artificial Sequence

<223> Trans-sialidase gene primer

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34

THE